

SEQUENCE LISTING

<110> Lovenberg, Timothy
Liu, Changlu

<120> DNAs Encoding Mammalian Histamine Receptor Of The H4 Subtype

<130> PRD-0033

<150> 09/790,849
<151> 2001-02-22

<150> 60/208,260
<151> 2000-05-31

<160> 27

<170> PatentIn version 3.2

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<212> DNA
<213> Homo sapiens

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<213> Homo sapiens

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Asn Ala Leu Val Ile Leu Ala Phe Val Val Asp Lys Asn Leu Arg His
35 40 45

Arg Ser Ser Tyr Phe Phe Leu Asn Leu Ala Ile Ser Asp Phe Phe Val
50 55 60

Gly Val Ile Ser Ile Pro Leu Tyr Ile Pro His Thr Leu Phe Glu Trp
65 70 75 80

Asp Phe Gly Lys Glu Ile Cys Val Phe Trp Leu Thr Thr Asp Tyr Leu
85 90 95

Leu Cys Thr Ala Ser Val Tyr Asn Ile Val Leu Ile Ser Tyr Asp Arg
100 105 110

Tyr Leu Ser Val Ser Asn Ala Val Ser Tyr Arg Thr Gln His Thr Gly
115 120 125

Val Leu Lys Ile Val Thr Leu Met Val Ala Val Trp Val Leu Ala Phe
130 135 140

Leu Val Asn Gly Pro Met Ile Leu Val Ser Glu Ser Trp Lys Asp Glu
145 150 155 160

Gly Ser Glu Cys Glu Pro Gly Phe Phe Ser Glu Trp Tyr Ile Leu Ala
165 170 175

Ile Thr Ser Phe Leu Glu Phe Val Ile Pro Val Ile Leu Val Ala Tyr
180 185 190

Phe Asn Met Asn Ile Tyr Trp Ser Leu Trp Lys Arg Asp His Leu Ser
195 200 205

Arg Cys Gln Ser His Pro Gly Leu Thr Ala Val Ser Ser Asn Ile Cys
210 215 220

Gly His Ser Phe Arg Gly Arg Leu Ser Ser Arg Arg Ser Leu Ser Ala
225 230 235 240

Ser Thr Glu Val Pro Ala Ser Phe His Ser Glu Arg Gln Arg Arg Lys
245 250 255

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260 265 270

Ala Ser Lys Met Gly Ser Phe Ser Gln Ser Asp Ser Val Ala Leu His

275

280

285

Gln Arg Glu His Val Glu Leu Leu Arg Ala Arg Arg Leu Ala Lys Ser
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Leu Ala Ile Leu Leu Gly Val Phe Ala Val Cys Trp Ala Pro Tyr Ser
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Leu Phe Thr Ile Val Leu Ser Phe Tyr Ser Ser Ala Thr Gly Pro Lys
 325 330 335

Ser Val Trp Tyr Arg Ile Ala Phe Trp Leu Gln Trp Phe Asn Ser Phe
 340 345 350

Val Asn Pro Leu Leu Tyr Pro Leu Cys His Lys Arg Phe Gln Lys Ala
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 <213> Mus musculus

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tggtacagca tagccttttg gctacagtgg ttcaattcac ttattaatcc ctttctatac	1080
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gtggacagaa atcttagaca tcgaagtaat tacttttttc ttaacttggc cattgcagac	180
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Asn	Ala	Val	Val	Ile	Leu	Ala	Phe	Val	Val	Asp	Arg	Asn	Leu	Arg	His
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 Gly Leu Ile Ser Ile Pro Leu Tyr Ile Pro His Val Leu Phe Asn Trp
 65 70 75 80
 Asn Phe Gly Ser Gly Ile Cys Met Phe Trp Leu Ile Thr Asp Tyr Leu
 85 90 95
 Leu Cys Thr Ala Ser Val Tyr Asn Ile Val Leu Ile Ser Tyr Asp Arg
 100 105 110
 Tyr Gln Ser Val Ser Asn Ala Val Ser Tyr Arg Ala Gln His Thr Gly
 115 120 125
 Ile Met Lys Ile Val Ala Gln Met Val Ala Val Trp Ile Leu Ala Phe
 130 135 140
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 145 150 155 160
 Thr Asn Thr Lys Asp Cys Glu Pro Gly Phe Val Thr Glu Trp Tyr Ile
 165 170 175
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 180 185 190
 Ala Tyr Phe Asn Val Gln Ile Tyr Trp Ser Leu Trp Lys Arg Arg Ala
 195 200 205
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 Ala Ser Gly His Leu His Arg Ala Gly Val Ala Cys Arg Thr Ser Asn
 225 230 235 240
 Pro Gly Leu Lys Glu Ser Ala Ala Ser Arg His Ser Glu Ser Pro Arg
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 Arg Lys Ser Ser Ile Leu Val Ser Leu Arg Thr His Met Asn Ser Ser
 260 265 270
 Ile Thr Ala Phe Lys Val Gly Ser Phe Trp Arg Ser Glu Ser Ala Ala
 275 280 285
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 Tyr Cys Leu Phe Thr Ile Val Leu Ser Thr Tyr Pro Arg Thr Glu Arg

325

330

335

Pro Lys Ser Val Trp Tyr Ser Ile Ala Phe Trp Leu Gln Trp Phe Asn
 340 345 350

Ser Phe Val Asn Pro Phe Leu Tyr Pro Leu Cys His Arg Arg Phe Gln
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 370 375 380

Gln Asn Gln Ser Val Ser Ser
 385 390

<210> 9
 <211> 391
 <212> PRT
 <213> Rattus rattus

<400> 9

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 35 40 45

Arg Ser Asn Tyr Phe Phe Leu Asn Leu Ala Ile Ser Asp Phe Phe Val
 50 55 60

Gly Val Ile Ser Ile Pro Leu Tyr Ile Pro His Thr Leu Phe Asn Trp
 65 70 75 80

Asn Pro Gly Ser Gly Ile Cys Met Phe Trp Leu Ile Thr Asp Tyr Leu
 85 90 95

Leu Cys Thr Ala Ser Val Tyr Ser Ile Val Leu Ile Ser Tyr Asp Arg
 100 105 110

Tyr Gln Ser Val Ser Asn Ala Val Arg Tyr Arg Ala Gln His Thr Gly
 115 120 125

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 130 135 140

Leu Val Asn Gly Pro Met Ile Leu Ala Ser Asp Ser Trp Lys Asn Ser
 145 150 155 160

Thr Asn Thr Glu Glu Cys Glu Pro Gly Phe Val Thr Glu Trp Tyr Ile
 165 170 175

Leu Ala Ile Thr Ala Phe Leu Glu Phe Leu Leu Pro Val Ser Leu Val

180

185

190

Val Tyr Phe Ser Val Gln Ile Tyr Trp Ser Leu Trp Lys Arg Gly Ser
195 200 205

Leu Ser Arg Cys Pro Ser His Ala Gly Phe Ile Ala Thr Ser Ser Arg
210 215 220

Gly Thr Gly His Ser Arg Arg Thr Gly Leu Ala Cys Arg Thr Ser Leu
225 230 235 240

Pro Gly Leu Lys Glu Pro Ala Ala Ser Leu His Ser Glu Ser Pro Arg
245 250 255

Gly Lys Ser Ser Leu Leu Val Ser Leu Arg Thr His Met Ser Gly Ser
260 265 270

Ile Ile Ala Phe Lys Val Gly Ser Phe Cys Arg Ser Glu Ser Pro Val
275 280 285

Leu His Gln Arg Glu His Val Glu Leu Leu Arg Gly Arg Lys Leu Ala
290 295 300

Arg Ser Leu Ala Val Leu Leu Ser Ala Phe Ala Ile Cys Trp Ala Pro
305 310 315 320

Tyr Cys Leu Phe Thr Ile Val Leu Ser Thr Tyr Arg Arg Gly Glu Arg
325 330 335

Pro Lys Ser Ile Trp Tyr Ser Ile Ala Phe Trp Leu Gln Trp Phe Asn
340 345 350

Ser Leu Ile Asn Pro Phe Leu Tyr Pro Leu Cys His Arg Arg Phe Gln
355 360 365

Lys Ala Phe Trp Lys Ile Leu Cys Val Thr Lys Gln Pro Ala Pro Ser
370 375 380

Gln Thr Gln Ser Val Ser Ser
385 390

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<212> PRT
<213> *Cavia porcellus*

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Val Val Val Ile Leu Ala Phe Ile Val Asp Arg Asn Leu Arg His Arg

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Ala	Ile	Ala	Ile	Pro	Leu	Tyr	Ile	Pro	Ser	Ser	Leu	Thr	Tyr	Trp	Thr
65					70					75					80
Ser	Gly	Lys	Gln	Ala	Cys	Val	Phe	Trp	Leu	Ile	Thr	Asp	Tyr	Leu	Leu
			85						90					95	
Cys	Thr	Ala	Ser	Val	Tyr	Asn	Ile	Val	Leu	Ile	Ser	Tyr	Asp	Arg	Tyr
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Gln	Ser	Val	Ser	Asn	Ala	Val	Trp	Tyr	Arg	Ala	Gln	His	Ser	Gly	Thr
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Trp	Lys	Ile	Ala	Thr	Gln	Met	Val	Ala	Val	Trp	Ile	Phe	Ser	Phe	Met
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Thr	Asn	Gly	Pro	Met	Ile	Leu	Ile	Ser	Asp	Ser	Trp	Gln	Asn	Ser	Thr
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Thr	Glu	Cys	Glu	Pro	Gly	Phe	Leu	Lys	Lys	Trp	Tyr	Phe	Ala	Leu	Pro
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Thr	Ser	Leu	Leu	Glu	Phe	Leu	Ile	Pro	Ile	Leu	Leu	Val	Ala	Tyr	Phe
			180					185					190		
Ser	Ala	His	Ile	Tyr	Trp	Ser	Leu	Trp	Lys	Arg	Glu	Lys	Leu	Ser	Arg
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Cys	Leu	Ser	His	Pro	Val	Leu	Pro	Ser	Asp	Ser	Ser	Ser	Ser	Asp	His
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225					230					235					240
Arg	Lys	Glu	Thr	Thr	Ala	Ser	Leu	Gly	Ser	Asp	Lys	Ser	Arg	Arg	Lys
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Ser	Ser	Leu	Leu	Pro	Ser	Ile	Arg	Ala	Tyr	Lys	Asn	Ser	Asn	Val	Ile
			260					265					270		
Ala	Ser	Lys	Met	Gly	Phe	Leu	Ser	His	Ser	Asp	Ser	Leu	Ala	Leu	Gln
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Gln	Arg	Glu	His	Ile	Glu	Leu	Phe	Arg	Ala	Arg	Lys	Leu	Ala	Lys	Ser
	290					295					300				
Leu	Ala	Ile	Leu	Leu	Ala	Ala	Phe	Ala	Ile	Cys	Trp	Ala	Pro	Tyr	Ser
305					310					315					320

Leu Thr Thr Val Ile Tyr Ser Phe Phe Pro Glu Arg Asn Leu Thr Lys
 325 330 335

Ser Thr Trp Tyr His Thr Ala Phe Trp Leu Gln Trp Phe Asn Ser Phe
 340 345 350

Val Asn Pro Phe Leu Tyr Pro Leu Cys His Lys Arg Phe Gln Lys Ala
 355 360 365

Phe Leu Lys Ile Leu Pro Val Arg Arg Gln Ser Thr Pro Pro His Asn
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Arg Ser Ile Ser Thr
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<210> 13
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<213> Artificial Sequence
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 <212> DNA
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<220>
<223> Oligonucleotide Primer

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